

Serial Number: 09/663,600A

ENTERED

#8

CNA

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patent bug). Sequences corrected: _____
- ☐ Other: _____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING

DATE: 10/30/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

PS

4 <110> APPLICANT: Dumas Milne Edwards, Jean-Baptiste
 5 Duclert, Aymeric
 6 Bougueleret, Lydie
 8 <120> TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS
 10 <130> FILE REFERENCE: 31.US3.CIP
 12 <140> CURRENT APPLICATION NUMBER: 09/663,600A
 13 <141> CURRENT FILING DATE: 2000-09-15
 15 <150> PRIOR APPLICATION NUMBER: 09/191,997
 16 <151> PRIOR FILING DATE: 1998-11-13
 18 <150> PRIOR APPLICATION NUMBER: 60/066,677
 19 <151> PRIOR FILING DATE: 1997-11-13
 21 <150> PRIOR APPLICATION NUMBER: 60/069,957
 22 <151> PRIOR FILING DATE: 1997-12-17
 24 <150> PRIOR APPLICATION NUMBER: 60/074,121
 25 <151> PRIOR FILING DATE: 1998-02-09
 27 <150> PRIOR APPLICATION NUMBER: 60/081,563
 28 <151> PRIOR FILING DATE: 1998-04-13
 30 <150> PRIOR APPLICATION NUMBER: 60/096,116
 31 <151> PRIOR FILING DATE: 1998-08-10
 33 <150> PRIOR APPLICATION NUMBER: 60/099,273
 34 <151> PRIOR FILING DATE: 1998-09-04
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 48 <222> LOCATION: 1
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 62 <210> SEQ ID NO: 3
 63 <211> LENGTH: 25
 64 <212> TYPE: DNA
 65 <213> ORGANISM: Artificial Sequence

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TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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67 <223> OTHER INFORMATION: Derivatized oligonucleotide for linking to mRNA

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69 atcaagaatt cgcacgagac catta 25
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73 <212> TYPE: DNA
74 <213> ORGANISM: Artificial Sequence

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82 <212> TYPE: DNA
83 <213> ORGANISM: Artificial Sequence

W--> 84 <220> FEATURE:
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87 ccgacaagac caacgtcaag gccgc 25
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90 <211> LENGTH: 25
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence

W--> 93 <220> FEATURE:
94 <223> OTHER INFORMATION: Alpha globin gene primer GLO-As

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99 <211> LENGTH: 25
100 <212> TYPE: DNA
101 <213> ORGANISM: Artificial Sequence

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103 <223> OTHER INFORMATION: Dehydrogenase gene primer 3 DH-S

W--> 104 <400> SEQUENCE: 7
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107 <210> SEQ ID NO: 8
108 <211> LENGTH: 25
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence

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112 <223> OTHER INFORMATION: Dehydrogenase gene primer 3 DH-As

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118 <212> TYPE: DNA
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001

TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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130 <223> OTHER INFORMATION: PP15 gene primer PP15-As
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134 <210> SEQ ID NO: 11
135 <211> LENGTH: 25
136 <212> TYPE: DNA
137 <213> ORGANISM: Artificial Sequence
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139 <223> OTHER INFORMATION: Elongation Factor A4 gene primer EFA1-S
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143 <210> SEQ ID NO: 12
144 <211> LENGTH: 25
145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
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148 <223> OTHER INFORMATION: Elongation Factor A4 gene primer EF1A-As
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152 <210> SEQ ID NO: 13
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154 <212> TYPE: DNA
155 <213> ORGANISM: Artificial Sequence
W--> 156 <220> FEATURE:
157 <223> OTHER INFORMATION: Primer derived from oligonucleotide of SEQ ID NO:3
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159 atcaagaatt cgcacgagac catta 25
161 <210> SEQ ID NO: 14
162 <211> LENGTH: 67
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
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166 <223> OTHER INFORMATION: PolydT primer
W--> 167 <220> FEATURE:
168 <221> NAME/KEY: misc_feature
169 <222> LOCATION: 67
170 <223> OTHER INFORMATION: n=a, g, c or t
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W--> 173 tttttvn 67
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RAW SEQUENCE LISTING

DATE: 10/30/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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184 <210> SEQ ID NO: 16
185 <211> LENGTH: 25
186 <212> TYPE: DNA
187 <213> ORGANISM: Artificial Sequence
W--> 188 <220> FEATURE:
189 <223> OTHER INFORMATION: Nested 3' primer
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191 cacgagagag actacacggt actgg                25
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194 <211> LENGTH: 526
195 <212> TYPE: DNA
196 <213> ORGANISM: Homo Sapiens
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198 <221> NAME/KEY: misc_feature
199 <222> LOCATION: complement(261..376)
200 <223> OTHER INFORMATION: blastn
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202 <221> NAME/KEY: misc_feature
203 <222> LOCATION: complement(380..486)
204 <223> OTHER INFORMATION: blastn
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206 <221> NAME/KEY: misc_feature
207 <222> LOCATION: complement(110..145)
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212 <223> OTHER INFORMATION: blastn
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214 <221> NAME/KEY: sig_peptide
215 <222> LOCATION: 90..140
216 <223> OTHER INFORMATION: Von Heijne matrix
W--> 217 <220> FEATURE:
218 <221> NAME/KEY: misc_feature
219 <222> LOCATION: 290
220 <223> OTHER INFORMATION: n=a, g, c or t
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222 aatatrarac agctacaata ttccagggcc artcacttgc catttctcat aacagcgtca    60
223 gagagaaaga actgactgar acgtttgag atg aag aaa gtt ctc ctc ctg atc    113
224                                     Met Lys Lys Val Leu Leu Leu Ile
225                                     -15                               -10
226 aca gcc atc ttg gca gtg gct gtw ggt ttc cca gtc tct caa gac cag    161

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RAW SEQUENCE LISTING

DATE: 10/30/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:35:00

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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227 Thr Ala Ile Leu Ala Val Ala Val Gly Phe Pro Val Ser Gln Asp Gln
228          -5                      1                      5
229 gaa cga gaa aaa aga agt atc agt gac agc gat gaa tta gct tca ggr      209
230 Glu Arg Glu Lys Arg Ser Ile Ser Asp Ser Asp Glu Leu Ala Ser Gly
231          10                      15                      20
232 wtt ttt gtg ttc cct tac cca tat cca ttt cgc cca ctt cca cca att      257
W--> 233 Xaa Phe Val Phe Pro Tyr Pro Tyr Pro Phe Arg Pro Leu Pro Pro Ile
234          25                      30                      35
W--> 235 cca ttt cca aga ttt cca tgg ttt aga cgt aan ttt cct att cca ata      305
W--> 236 Pro Phe Pro Arg Phe Pro Trp Phe Arg Arg Xaa Phe Pro Ile Pro Ile
237 40                      45                      50                      55
238 cct gaa tct gcc cct aca act ccc ctt cct agc gaa aag taaacaaraa      354
239 Pro Glu Ser Ala Pro Thr Thr Pro Leu Pro Ser Glu Lys
240          60                      65
241 ggaaaagtca crataaacct ggtcacctga aattgaaatt gagccacttc cttgaaraat      414
242 caaaaattcct gttataaaaa raaaaacaaa tgtaattgaa atagcacaca gcattctcta      474
243 gtcaatatct ttagtgatct tctttaataa acatgaaagc aaaaaaaaaa aa      526
245 <210> SEQ ID NO: 18
246 <211> LENGTH: 17
247 <212> TYPE: PRT
248 <213> ORGANISM: Homo Sapiens
W--> 249 <220> FEATURE:
250 <221> NAME/KEY: SIGNAL
251 <222> LOCATION: 1..17
252 <223> OTHER INFORMATION: Von Heijne matrix
253     score 8.2
254     seq LLLITAILAVAVG/FP
W--> 255 <400> SEQUENCE: 18
256 Met Lys Lys Val Leu Leu Leu Ile Thr Ala Ile Leu Ala Val Ala Val
257 1                      5                      10                      15
258 Gly
260 <210> SEQ ID NO: 19
261 <211> LENGTH: 822
262 <212> TYPE: DNA
263 <213> ORGANISM: Homo Sapiens
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266 <222> LOCATION: 260..464
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270 <222> LOCATION: 118..184
271 <223> OTHER INFORMATION: blastn
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273 <221> NAME/KEY: misc_feature
274 <222> LOCATION: 56..113
275 <223> OTHER INFORMATION: blastn
W--> 276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature

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PSI Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001

TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

L:44 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <400> field identifier
L:57 M:283 W: Missing Blank Line separator, <220> field identifier
L:59 M:283 W: Missing Blank Line separator, <400> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:75 M:283 W: Missing Blank Line separator, <220> field identifier
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:84 M:283 W: Missing Blank Line separator, <220> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:93 M:283 W: Missing Blank Line separator, <220> field identifier
L:95 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:120 M:283 W: Missing Blank Line separator, <220> field identifier
L:122 M:283 W: Missing Blank Line separator, <400> field identifier
L:129 M:283 W: Missing Blank Line separator, <220> field identifier
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L:167 M:283 W: Missing Blank Line separator, <220> field identifier
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L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
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L:201 M:283 W: Missing Blank Line separator, <220> field identifier
L:205 M:283 W: Missing Blank Line separator, <220> field identifier
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L:217 M:283 W: Missing Blank Line separator, <220> field identifier
L:221 M:283 W: Missing Blank Line separator, <400> field identifier
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L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:249 M:283 W: Missing Blank Line separator, <220> field identifier
L:255 M:283 W: Missing Blank Line separator, <400> field identifier
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VERIFICATION SUMMARY

DATE: 10/30/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:623 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31
L:810 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:34
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L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:964 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:1272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
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L:2600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
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L:3154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
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L:3403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3407 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56

VERIFICATION SUMMARY

DATE: 10/30/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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L:3847 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59

1600

RAW SEQUENCE LISTING

DATE: 10/17/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:26:46

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Output Set: N:\CRF3\10172001\I663600A.raw

4 <110> APPLICANT: Dumas Milne Edwards, Jean-Baptiste
 5 Duclert, Aymeric
 6 Bougueleret, Lydie
 8 <120> TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS
 10 <130> FILE REFERENCE: 31.US3.CIP
 12 <140> CURRENT APPLICATION NUMBER: 09/663,600A
 13 <141> CURRENT FILING DATE: 2000-09-15
 15 <150> PRIOR APPLICATION NUMBER: 09/191,997
 16 <151> PRIOR FILING DATE: 1998-11-13
 18 <150> PRIOR APPLICATION NUMBER: 60/066,677
 19 <151> PRIOR FILING DATE: 1997-11-13
 21 <150> PRIOR APPLICATION NUMBER: 60/069,957
 22 <151> PRIOR FILING DATE: 1997-12-17
 24 <150> PRIOR APPLICATION NUMBER: 60/074,121
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 27 <150> PRIOR APPLICATION NUMBER: 60/081,563
 28 <151> PRIOR FILING DATE: 1998-04-13
 30 <150> PRIOR APPLICATION NUMBER: 60/096,116
 31 <151> PRIOR FILING DATE: 1998-08-10
 33 <150> PRIOR APPLICATION NUMBER: 60/099,273
 34 <151> PRIOR FILING DATE: 1998-09-04
 36 <160> NUMBER OF SEQ ID NOS: 229
 38 <170> SOFTWARE: Patent.pm

P. 2
 Does Not Comply
 Corrected Diskette Needed

ERRORED SEQUENCES

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12237 <211> LENGTH: 142

12238 <212> TYPE: PRT

12239 <213> ORGANISM: Homo sapiens

12240 <400> SEQUENCE: 229

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 12246 Ile Met Lys Val Asp Lys Asp Arg Gln Met Val Val Leu Glu Glu Glu
 12247 35 40 45
 12248 Phe Arg Asn Ile Ser Pro Glu Glu Leu Lys Met Glu Leu Pro Glu Arg
 12249 50 55 60
 12250 Gln Pro Arg Phe Val Val Tyr Ser Tyr Lys Tyr Val Arg Asp Asp Gly
 12251 65 70 75 80
 12252 Arg Val Ser Tyr Pro Leu Cys Phe Ile Phe Ser Ser Pro Val Gly Cys
 12253 85 90 95
 12254 Lys Pro Glu Gln Gln Met Met Tyr Ala Gly Ser Lys Asn Arg Leu Val
 12255 100 105 110
 12256 Gln Thr Ala Glu Leu Thr Lys Val Phe Glu Ile Arg Thr Thr Asp Asp

RAW SEQUENCE LISTING

DATE: 10/17/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:26:49

Input Set : D:\Seqlist.txt

Output Set: N:\CRF3\10172001\I663600A.raw

12257 115 120 125
12258 Leu Thr Glu Ala Trp Leu Gln Glu Lys Leu Ser Phe Phe Arg
12259 130 135 140
E--> 12260 1

Errored Delete End of File Non ASCII Text

VERIFICATION SUMMARY

DATE: 10/17/2001

PATENT APPLICATION: US/09/663,600A

TIME: 12:26:50

Input Set : D:\Seqlist.txt

Output Set: N:\CRF3\10172001\I663600A.raw

L:44 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <400> field identifier
L:57 M:283 W: Missing Blank Line separator, <220> field identifier
L:59 M:283 W: Missing Blank Line separator, <400> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:75 M:283 W: Missing Blank Line separator, <220> field identifier
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:84 M:283 W: Missing Blank Line separator, <220> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:93 M:283 W: Missing Blank Line separator, <220> field identifier
L:95 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:120 M:283 W: Missing Blank Line separator, <220> field identifier
L:122 M:283 W: Missing Blank Line separator, <400> field identifier
L:129 M:283 W: Missing Blank Line separator, <220> field identifier
L:131 M:283 W: Missing Blank Line separator, <400> field identifier
L:138 M:283 W: Missing Blank Line separator, <220> field identifier
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
L:147 M:283 W: Missing Blank Line separator, <220> field identifier
L:149 M:283 W: Missing Blank Line separator, <400> field identifier
L:156 M:283 W: Missing Blank Line separator, <220> field identifier
L:158 M:283 W: Missing Blank Line separator, <400> field identifier
L:165 M:283 W: Missing Blank Line separator, <220> field identifier
L:167 M:283 W: Missing Blank Line separator, <220> field identifier
L:171 M:283 W: Missing Blank Line separator, <400> field identifier
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:179 M:283 W: Missing Blank Line separator, <220> field identifier
L:181 M:283 W: Missing Blank Line separator, <400> field identifier
L:188 M:283 W: Missing Blank Line separator, <220> field identifier
L:190 M:283 W: Missing Blank Line separator, <400> field identifier
L:197 M:283 W: Missing Blank Line separator, <220> field identifier
L:201 M:283 W: Missing Blank Line separator, <220> field identifier
L:205 M:283 W: Missing Blank Line separator, <220> field identifier
L:209 M:283 W: Missing Blank Line separator, <220> field identifier
L:213 M:283 W: Missing Blank Line separator, <220> field identifier
L:217 M:283 W: Missing Blank Line separator, <220> field identifier
L:221 M:283 W: Missing Blank Line separator, <400> field identifier
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:249 M:283 W: Missing Blank Line separator, <220> field identifier
L:255 M:283 W: Missing Blank Line separator, <400> field identifier
L:264 M:283 W: Missing Blank Line separator, <220> field identifier

VERIFICATION SUMMARY

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L:268 M:283 W: Missing Blank Line separator, <220> field identifier
L:272 M:283 W: Missing Blank Line separator, <220> field identifier
L:276 M:283 W: Missing Blank Line separator, <220> field identifier
L:280 M:283 W: Missing Blank Line separator, <220> field identifier
L:284 M:283 W: Missing Blank Line separator, <220> field identifier
L:288 M:283 W: Missing Blank Line separator, <220> field identifier
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:623 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31 *ok*
L:810 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:34
L:927 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:964 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:1272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:1492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1804 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1810 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:2351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:2496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:2926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2950 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:3104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:3154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:3401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3407 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56

VERIFICATION SUMMARY

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Input Set : D:\Seqlist.txt

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L:3651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:3660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:3838 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3847 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:12260 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:229 @✓